-continued

- 1. A chimeric single guide RNA molecule (sgRNA), comprising:
 - (i) a guide sequence capable of hybridizing to a target sequence in a genomic locus of interest in a eukaryotic cell, wherein the guide sequence has a length of 21 to 23 nucleotides and comprises a 5'-terminal guanosine
 (G) residue,
- (ii) a tracr-mate sequence capable of hybridizing with a tracr sequence, and
- (iii) a tracr sequence;
- wherein the sgRNA is capable of forming a CRISPR-Cas complex with *Staphylococcus aureus* Cas9 (SaCas9).
- 2. The sgRNA of claim 1, wherein a poly U tract in wild-type tracr-mate sequence, which is located at positions 2-5 starting from 3'-end of the guide sequence, is replaced with poly C.
- **3**. The sgRNA of claim **2**, wherein a poly A tract in wild-type tracr sequence, which basepairs with the poly U tract in wild-type tracr-mate sequence, is replaced with poly G
- **4**. The sgRNA of claim **1**, wherein the guide sequence has a length of 21 nucleotides.
- 5. The sgRNA of claim 1, wherein the guide sequence has a length of 22 nucleotides.
- **6**. The sgRNA of claim **1**, wherein the guide sequence has a length of 23 nucleotides.
- 7. The sgRNA of claim 1, wherein a repeat:anti-repeat duplex formed by hybridization of the tracr-mate sequence and the tracr sequence is truncated compared to that of a wild-type *Staphylococcus aureus* CRISPR-Cas9 system.
 - **8-12**. (canceled)
- 13. The sgRNA of claim 7, wherein the tracr-mate sequence has a length of 14-25 nucleotides.
 - 14-17. (canceled)

- 18. The sgRNA of claim 1, wherein the tracr sequence has a length of 50-100 nucleotides.
- 19. The sgRNA of claim 1, wherein the sgRNA comprises at least one loop that comprises an insertion of an RNA sequence capable of binding to an adaptor protein.
- 20. The sgRNA of claim 19, wherein the inserted RNA sequence is an aptamer.
- 21. A composition comprising a CRISPR-Cas complex, wherein the CRISPR-Cas complex comprises the sgRNA of claim 1 and a SaCas9 protein.
- 22. The composition of claim 21, wherein the SaCas9 protein is linked to a heterologous functional domain, optionally wherein the heterologous functional domain is a transcriptional activation domain or a transcriptional repressor domain.
 - 23. (canceled)
- ${\bf 24}.$ A composition comprising the sgRNA of claim ${\bf 20}$ bound to an adaptor protein.
- 25. The composition of claim 24, wherein the adaptor protein is linked to a heterologous functional domain, optionally wherein the heterologous functional domain is a transcriptional activation domain or a transcriptional repressor domain.
 - 26. (canceled)
- 27. A polynucleotide encoding the sgRNA of claim 1.
- **28**. A vector comprising the polynucleotide of claim **27** operably linked to a promoter.
- **29**. The vector of claim **28**, further comprising a polynucleotide encoding a SaCas9 protein.
 - 30. A eukaryotic cell comprising the sgRNA of claim 1.
- 31. A eukaryotic cell transformed with the vector of claim 28.

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